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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/777,526A

DATE: 06/11/2002

TIME: 16:40:06

Input Set : A:\HYZ-030CPCN3(47508-518).ST25.txt  
Output Set: N:\CRF3\06112002\I777526A.rawDoes Not Comply  
Corrected Diskette Needed

3 <110> APPLICANT: AGRAWAL, Sudhir  
 4 DIASIO, Robert H.  
 5 ZHANG, Ruiwen  
 7 <120> TITLE OF INVENTION: A Method of Down-Regulating Gene Expression  
 9 <130> FILE REFERENCE: HYZ-030CPCN3 (47508-518)  
 11 <140> CURRENT APPLICATION NUMBER: US 09/777,526A  
 C--> 12 <141> CURRENT FILING DATE: 2002-06-04  
 14 <150> PRIOR APPLICATION NUMBER: US 08/758,005  
 15 <151> PRIOR FILING DATE: 1996-11-27  
 17 <150> PRIOR APPLICATION NUMBER: US 08/709,910  
 18 <151> PRIOR FILING DATE: 1996-09-09  
 20 <150> PRIOR APPLICATION NUMBER: US 08/328,520  
 21 <151> PRIOR FILING DATE: 1994-10-25  
 23 <160> NUMBER OF SEQ ID NOS: 21  
 25 <170> SOFTWARE: PatentIn version 3.0

## ERRORED SEQUENCES

300 <210> SEQ ID NO: 21  
 301 <211> LENGTH: 18  
 302 <212> TYPE: DNA  
 303 <213> ORGANISM: Artificial Sequence  
 305 <220> FEATURE:  
 306 <223> OTHER INFORMATION: Mismatched Control  
 307 DNA/RNA  
 310 <400> SEQUENCE: 21  
 311 gcaugcatcc gcacaggc  
 E--> 313 hyz-030cpcn3(47508-518).st25  
 E--> 315 Page 6

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Delete

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/777,526A

DATE: 06/11/2002

TIME: 16:40:07

Input Set : A:\HYZ-030CPCN3(47508-518).ST25.txt  
Output Set: N:\CRF3\06112002\I777526A.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:313 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:21  
L:313 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21  
L:313 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:18  
L:313 M:254 E: No. of Bases conflict, LENGTH:Input:25 Counted:32 SEQ:21  
L:313 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:19  
L:313 M:112 C: (48) String data converted to lower case,  
M:254 Repeated in SeqNo=21  
L:315 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2  
L:315 M:252 E: No. of Seq. differs, <211> LENGTH:Input:18 Found:36 SEQ:21